

STOMACH CONTENTS OF LONG-FINNED PILOT WHALES (*GLOBICEPHALA MELAS*) STRANDED ON THE U.S. MID-ATLANTIC COAST

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ABSTRACT

Ten prey taxa were recorded from the stomach contents of eight long-finned pilot whales (*Globicephala melas*) independently stranded along the U.S. mid-Atlantic coast. Relative importance of prey species was determined by methods that incorporate prey frequencies of occurrence, proportions of numerical abundance, and proportions of reconstructed mass. Separate analyses of trace (free, durable body parts representing well-digested prey items) and non-trace (relatively intact prey specimens) food material were conducted in order to address biases caused by differential rates of digestion and passage through the gastrointestinal tract. Different measures of prey importance yielded varying results, but the long-finned squid (*Loligo pealei*) was the most important prey species regardless of how prey importance was defined. Fishes were relatively unimportant in the diet. Our results indicate that the diets of western North Atlantic long-finned pilot whales differ substantially from what has been previously reported in the literature and that results from food-habits studies that utilize different techniques may not be comparable.

Key words: *Globicephala melas*, long-finned pilot whale, stomach contents, food habits, stranding, *Loligo pealei*, prey importance.

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Whales of the genus *Globicephala*, including long-finned (*G. melas*) and short-finned (*G. macrorhynchus*) pilot whales, are thought to account for a large portion of the food energy consumed by cetaceans on the outer continental shelf of the northeastern United States north of Cape Hatteras (Kenney *et al.* 1985). *G. melas* has a more northern distribution than *G. macrorhynchus* (Winn 1982, Payne and Heinemann 1993) and is probably the most abundant of the two pilot whale species in this region (Payne and Heinemann 1993).

Our motivation for undertaking this project came from the realization that previous studies of *G. melas* food habits in the western North Atlantic have been characterized by low prey diversities relative to studies conducted in the eastern North Atlantic and the Southern Hemisphere (Table 1; hereafter, "pilot whale" refers to *G. melas*). We wanted to determine whether this difference was real or merely an artifact of differences in research methodologies used in the three areas. Two major pilot whale food-habits studies have been conducted in the western North Atlantic: Sergeant (1962) and Overholtz and Waring (1991). Using stomach contents data from animals taken in the Newfoundland drive fishery, Sergeant (1962) found that pilot whales primarily fed on the short-finned squid (*Illex illecebrosus*) and supplemented their diet with cod (*Gadus morhua*) when squid were scarce. Overholtz and Waring (1991) concluded from their study of five pilot whales incidentally captured by the pelagic mackerel fishery that Atlantic mackerel (*Scomber scombrus*) represented 71% of the diet and the remaining 29% consisted of long-finned squid (*Loligo pealei*) by existing mass.

In addition to these two food-habits studies in the western North Atlantic, three other papers contain brief information on stomach contents. Mercer (1967) reported that the stomachs of two pilot whales from an anomalous pod that overwintered in Newfoundland waters contained only Greenland turbot (*Reinhardtius hippoglossoides*). In a more recent report on the ecological relationship of pilot whales and *I. illecebrosus*, Mercer (1975) cited an observation of the squid *Gonatus fabricii* in the stomachs of 12 whales driven ashore in Newfoundland. In a paper on the incidental catch of marine mammals in foreign fisheries off the northeast United States, Waring *et al.* (1990) stated

Table 1. Number of prey species of long-finned pilot whales reported in the literature for three geographical regions (n = the number of stomachs examined).

	NW Atlantic ¹ $n > 51$	NE Atlantic ² $n > 861$	Tasmania ³ $n = 2$
Cephalopoda	3	15	14
Fishes	4	16	1
Crustacea	0	2	0
Others	0	4	0
TOTAL	7	37	15

From: Jensen 1916², Ritchie 1924², Sergeant 1962¹, Mercer 1967¹, Mercer 1975¹, Martin *et al.* 1987², Waring *et al.* 1990¹, Overholtz and Waring 1991¹, Gales and Pemberton 1992³, Desportes and Mouritsen 1993².

that the stomachs of two pilot whales captured by the mackerel fishery contained only mackerel.

It should be noted that all of the studies in the western North Atlantic relied heavily on intact prey items to determine relative dietary importance, while work in other areas (e.g., Martin *et al.* 1987, Gales and Pemberton 1992, Desportes and Mouritsen 1993) examined both intact and well-digested food material. Because of differences in digestion rates between fish and squid flesh (Bigg and Fawcett 1985), estimates of the dietary importance of cephalopods may be negatively biased when only intact prey are quantified or when existing prey mass is used.

Stomach contents of stranded animals may exhibit biases associated with the stranding event, but they do provide insight into dietary habits of the species. In this paper we describe the stomach contents of eight long-finned pilot whales that stranded independently along the mid-Atlantic coast of the United States. These eight stomachs represent the entire food-habits collection from stranded *G. melas* at the Smithsonian Institution's National Museum of Natural History. Several analytical methods were used to achieve a complete picture of the diet.

METHODS

Sample collection—Stranding locations of the eight whales are plotted in Figure 1. Table 2 provides information on stranding location, date, gender, body length, and maturity status for each animal. Based on body length, individuals were classified as calves (not fully weaned), immature (nutritionally independent but sexually immature), or sexually mature.

Stomachs were removed, after the esophagus and intestine had been ligated, then transported to the Smithsonian Institution where they were temporarily stored frozen. Stomach contents were subsequently placed in 70% ethanol for long-term storage at the Smithsonian. During examination, all prey items were passed through a metal sieve with a mesh diameter of 1.0 mm to isolate hard parts.

Prey identification—Prey items were identified by comparing whole specimens, and parts thereof, to a laboratory reference collection, museum reference collections at the Smithsonian and the Harvard University Museum of Comparative Zoology, and published guides, including Flescher 1980, Clarke 1986a, Vecchione *et al.* 1989, and Roper *et al.* 1984. Structures used to identify partially digested food items included cephalopod lower beaks, teleost sagittal otoliths, teleost dentaries, skulls, squalid spines, and articulated vertebral columns.

Prey importance—Separate analyses of "trace" and "non-trace" (Bigg and Perez 1985) prey categories were conducted. Trace food items were those that were well digested and represented only by disarticulated skeletal elements, squid beaks, or squalid spines. The non-trace category consisted of relatively undigested food items. Whole specimens, fish heads with attached flesh, torsos, cephalopod mantles, buccal masses with beaks, and crowns (arms with buccal

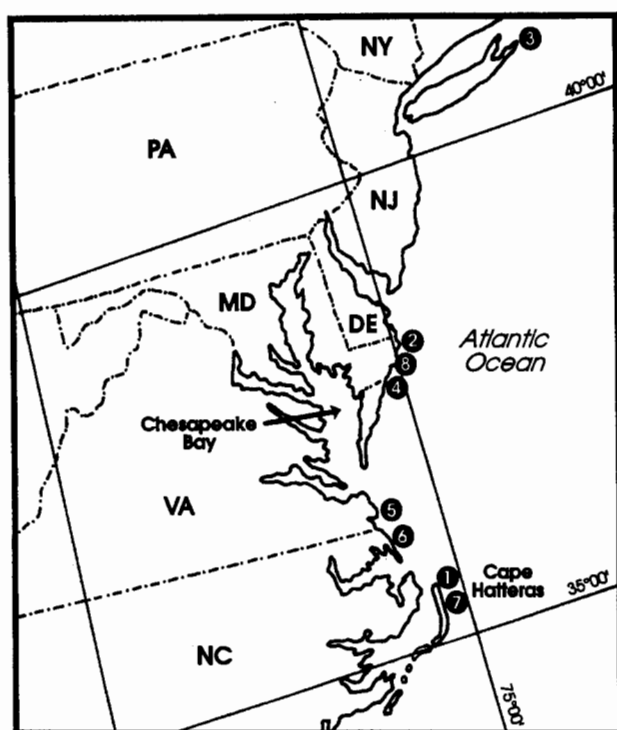


Figure 1. Stranding locations of long-finned pilot whales included in food-habits analyses.

Table 2. Summary of stranding data for long-finned pilot whales included in food-habits analyses (No. = identification number, Yr = Year, Mo = Month, D = Day, Len = Standard Length in Meters).

No. ¹	State	Yr	Mo	D	Len	Sex	Maturity ²
1	NC	73	3	29	2.15	F	Calf
2	MD	77	5	05	3.04	M	Immature
3	NY	79	3	27	2.80	M	Calf
4	VA	82	4	12	3.95	M	Immature
5	VA	87	3	17	5.05	F	Mature
6	NC	87	4	21	3.88	F	Mature
7	NC	88	9	28	2.35	M	Calf
8	MD	93	4	10	4.05	F	Mature

¹ Correspond to numbers in Figure 1.

² From morphometric data given by Sergeant 1962, Kasuya *et al.* 1988, Bloch *et al.* 1993, Desportes *et al.* 1993, Martin and Rothery 1993.

Table 3. Regression equations used to estimate length (dorsal mantle length for squids, fork length for teleosts, and total length for squalids: L) and weight (W) of pilot whale prey items from otolith length (OL), dentary length (DL), and cephalopod lower rostral length (LRL). All lengths are in millimeters and weights are in grams.

Prey species	Regression equation	Source
<i>Loligo pealei</i> (long-finned squid)	$\log L = 1.767 + 1.4 \log LRL$ $W = 0.25662(L/10)^{2.15182}$	This study Lange and Johnson 1981
<i>Ommastrephidae</i> ¹ (short-finned squid)	$\ln W = 1.773 + 2.40 \ln LRL$	Clarke 1962
<i>Histioteuthidae</i> ²	$L = -13.6 + 22.21 LRL$ $\ln W = 1.594 + 2.31 \ln LRL$	Clarke 1986a Clarke 1986a
<i>Chiroteuthis veranyi</i>	$\ln W = -0.241 + 2.7 \ln LRL$	Clarke 1980
<i>Squalus acanthias</i> (spiny dogfish)	$\ln[(W + 1)/1000] =$ $0.0293(L/10) - 1.1714$ females ≥ 500 mm $\ln[(W + 1)/1000] =$ $0.0235(L/10) - 0.8535$ males ≥ 500 mm	Nammack et al. 1985 Nammack et al. 1985
<i>Scomber scombrus</i> (Atlantic mackerel)	$L = -10.04 + 12.66 DL$ $L = 10(7.33 OL + 0.37)$ $W = 0.00756(L/10)^{3.082}$	This study Recchia and Read 1989 Kulka and Stobo 1981
<i>Clupea harengus</i> (Atlantic herring)	$L = 69.23 OL - 27.48$ $\log W = 3.12 \log L - 5.41$	Recchia and Read 1989 Recchia and Read 1989
<i>Merluccius bilinearis</i> (silver hake)	$L = 20.9 OL - 0.41$ $\log W = 3.08 \log(L/10) - 2.23$	Recchia and Read 1989 Kohler et al. 1970
<i>Urophycis</i> spp. (other hake)	$L = (1.52 OL)^{1.1456} 10$ $W = 0.003998(L/10)^{3.1718}$	Bowen and Harrison 1994 Bowen and Harrison 1994

¹ Most of the ommastrephids were believed to be *Illex illecebrosus*, so the regression equation used to calculate ommastrephid mass was that given by Clarke (1962) for *I. illecebrosus*.

² Regressions are available for the family Histioteuthidae, but not for the species *Histioteuthis reversa*.

mass and beak) all qualified as non-trace. Analysis of trace prey was performed in order to address biases caused by differential rates of digestion.

Size of prey at ingestion was reconstructed by estimating mass from body length or the length of undigested hard parts (Clarke 1962, Kohler et al. 1970, Clarke 1980, Lange and Johnson 1981, Nammack et al. 1985, Clarke 1986a, Recchia and Read 1989, Bowen and Harrison 1994). Table 3 lists the regression equations used to estimate size of prey from otolith lengths, dentary lengths, cephalopod lower beak rostral lengths, and body lengths. The regression equation relating dorsal mantle length and lower rostral length for *Loligo pealei* was calculated using a sample of 25 specimens ($r^2 = 0.728$). Mackerel otoliths were very fragile, making them unsuitable for use in body-

size estimation. Dentary bones were the most durable structures from mackerel, and we used these for identification and body-size estimation. Disarticulated dentaries were measured in a straight line from the anterior tip to the posterior margin, along the dorsal surface. The regression of fork length on dentary length for *Scomber scombrus* was estimated from a sample of 49 specimens ($r^2 = 0.993$). No length or weight regressions were available for *Seleneuthis scintillans* (of the family Lycoteuthidae), a small squid growing to a mantle length of about 30 mm (Clarke 1986a). The three *S. scintillans* beaks recovered were assumed to represent an insignificant portion of the overall diet and were excluded from further analysis. Sexual dimorphism necessitated that separate length/weight regressions be calculated for male and female spiny dogfish, *Squalus acanthias* (Nammack *et al.* 1985). It was impossible to determine the sex of these ingested prey, so the reconstructed mass of each dogfish was the average of the mass of a male and a female of the corresponding length.

Left and right otoliths, left and right dentaries, and upper and lower beaks were separated, whenever possible. The minimum number of individuals present from each species was determined by the largest number of component parts from paired structures. When digestion precluded distinction of right and left otoliths, the total number of damaged otoliths from each species was divided by two in order to estimate the number of individual fish. All otoliths, dentaries, and beaks were counted, but only intact structures were measured to find the average size of each individual. Otoliths were categorized on a scale from 0 (undamaged otoliths retrieved from skulls) to 5 (severely degraded free otoliths) following the methods of Recchia and Read (1989). Otoliths categorized as 3 or higher were not used in size estimations, unless no undamaged otoliths were present. When intact hard parts were unavailable to estimate body size for a prey species in a particular stomach, the available skeletal structures were measured, which may have resulted in an underestimation of the mass represented by those structures. No mackerel dentaries recovered appeared to be damaged. Few squid beaks were too damaged to estimate size.

The relative importance of prey was measured by nine methods: (1) trace frequency of occurrence, (2) non-trace frequency of occurrence, (3) trace numerical abundance, (4) non-trace numerical abundance, (5) trace proportion of mass, (6) non-trace proportion of mass, (7) trace index of relative importance, (8) non-trace index of relative importance, and (9) "modified mass." Frequency of occurrence (FO) is the proportion of stomachs that contained a particular prey category, regardless of its mass or abundance. The denominator used in the calculation of trace frequency of occurrence is equal to the total number of stomachs in the sample that contained trace food remains ($n = 8$), while the denominator in non-trace FO is the number of stomachs that contained non-trace prey remains ($n = 4$). Proportion of numerical abundance (%Num) is the percentage of the total number of prey items recovered from all stomachs represented by a particular prey category. Proportion of reconstructed mass (%Mass) is the percentage of mass of prey at ingestion recovered from all stomachs represented by a particular prey category. Ingested prey mass was

reconstructed to address biases caused by differential digestion rates. When diagnostic hard parts from more than 25 trace specimens of a prey taxon were present in one stomach, all hard parts from that taxon were tallied, and then a subsample of 25 was randomly selected and measured to estimate the average mass of individuals from that taxon in that particular stomach. Index of relative importance (IRI; Pinkas *et al.* 1971) combines the three methods already described and is calculated by the following equation:

$$\text{IRI} = \text{FO} \times (\% \text{Num} + \% \text{Mass})$$

The final method, modified mass, was adapted from the "modified volume" of Bigg and Perez (1985). It integrated frequency and mass measurements of both trace and non-trace food remains. Reconstructed mass was used rather than volume because we were interested in the sizes of prey at ingestion (and not the existing prey sizes after some unknown period of digestion, given the fact that different species digest at different rates). Prey items representing less than 1% of total mass were excluded from modified mass calculations (Bigg and Perez 1985). Modified mass was calculated by: (1) Determining the proportion of all fish to all squid by non-trace FO. (2) Determining the proportion of each species within these categories by total %Mass (trace and non-trace %Mass combined). (3) Adjusting the mass ratios for each species to sum to the total proportions of squid and fish present in the diet. (4). Readjusting all values to sum to 100%.

Some cephalopod beaks and teleost otoliths were not identified, either because they were not represented in reference collections or because they were severely damaged. All unidentified beaks were pooled and included in the calculations of trace frequency of occurrence and trace numerical abundance. Since the mass represented by these unidentified hard parts was unknown, they were excluded from reconstructed mass calculations.

Prey species compositions (measured by the above methods) were calculated for all eight whales as a group. Because of the small sample size, detailed investigations of potential dietary differences between age, sex, or reproductive categories were not possible.

RESULTS

Table 4 lists the ten prey species identified in the stomachs of stranded pilot whales. The long-finned squid, *Loligo pealei*, was the most important prey item, regardless of how prey importance was defined (Tables 5 through 7). Measures of prey importance for *L. pealei* ranged from 47.4% (modified mass) to 93.8% (non-trace numerical abundance) of the overall diet. Pilot whales fed on *L. pealei* ranging in size from 50 to 420 mm mantle length, but most were between 100 and 300 mm (Fig. 2). The largest *L. pealei* weighed approximately 798 g. *Histioteuthis reversa* and squids of the family Ommastrephidae consistently ranked in the top three prey taxa by trace measures (Table 5). Ommastrephids and *H. reversa* were present in 62.5% and 50% of stomachs, respectively, but were not represented by any non-trace specimens. Un-

Table 4. List of prey species present in the stomachs of long-finned pilot whales stranded along the U.S. mid-Atlantic coast, with their associated trace and non-trace abundances.

Species name	Common name	No. of trace specimens	No. of non-trace specimens
FISHES			
<i>Clupea harengus</i>	Atlantic herring	8	0
<i>Merluccius bilinearis</i>	Silver hake	4	0
<i>Scomber scombrus</i>	Atlantic mackerel	2	2
<i>Squalus acanthias</i>	Spiny dogfish	3	3
<i>Urophycis</i> spp.	Hake (red, white)	5	0
Unknown fish		1	0
SQUIDS			
<i>Chiroteuthis veranyi</i>		87	0
<i>Histioteuthis reversa</i>		221	0
<i>Loligo pealei</i>	Long-finned squid	560	75
Ommastrephidae	Short-finned squid	184	0
<i>Selenoteuthis scintillans</i>		3	0
Unknown squid		62	0

identified squid beaks were present in four stomachs and represented 5.1% of the trace numerical abundance.

Fish were relatively unimportant in the diet of these eight animals. Surprisingly, spiny dogfish, present only in one stomach, appeared to be the most important fish species. The contribution of dogfish to the overall diet ranged from 0.3% (trace numerical abundance) to 38.3% (non-trace proportion of

Table 5. Frequency of occurrence (FO), proportion of numerical abundance (%Num), and proportion of reconstructed mass (%Mass), with associated ranks (#), for the trace food materials of long-finned pilot whales stranded along the U.S. mid-Atlantic coast.

Species	Trace FO		Trace %Num		Trace %Mass	
	%	#	%	#	%	#
<i>L. pealei</i>	75.0%	1	49.1%	1	73.1%	1
Ommastrephidae	62.5%	2	16.1%	3	12.4%	2
<i>H. reversa</i>	50.0%	3	19.4%	2	7.1%	3
<i>S. acanthias</i>	12.5%	7	<1%	6	4.3%	4
<i>C. veranyi</i>	50.0%	3	7.6%	4	1.5%	5
<i>C. harengus</i>	12.5%	7	<1%	6	1.3%	6
<i>S. scombrus</i>	12.5%	7	<1%	6	<1%	7
<i>Urophycis</i> spp.	12.5%	7	<1%	6	<1%	7
<i>M. bilinearis</i>	12.5%	7	<1%	6	<1%	7
<i>S. scintillans</i>	37.5%	6	<1%	6	—	—
Unknown squid	50.0%	3	5.1%	5	—	—
Unknown fish	12.5%	7	<1%	6	—	—

Table 6. Frequency of occurrence (FO), proportion of numerical abundance (%Num), and proportion of reconstructed mass (%Mass), with associated ranks (#), for the non-trace food materials of long-finned pilot whales stranded along the U.S. mid-Atlantic coast.

Species	Non-trace FO		Non-trace %Num		Non-trace %Mass	
	%	#	%	#	%	#
<i>L. pealei</i>	75.0%	1	93.8%	1	51.9%	1
<i>S. acanthias</i>	25.0%	2	3.8%	2	38.3%	2
<i>S. scombrus</i>	25.0%	2	2.5%	3	9.8%	3

mass). Dogfish rivaled long-finned squid in non-trace mass (Table 6) and modified mass (Table 7), even though only six individuals were recovered (three intact and three trace). Each dogfish was about 750 mm in length and weighed about 1,640 g. Atlantic mackerel contributed a minimum of 0.2% to the overall diet when measured by either trace numerical abundance or trace proportion of mass, and a maximum of 25% when measured by non-trace frequency of occurrence. Unidentified otoliths were present in one stomach and represented 0.08% of the diet by trace numerical abundance. Dietary importance rankings of the three species represented by non-trace specimens were all in agreement, with long-finned squid first, spiny dogfish second, and Atlantic mackerel third (Table 6). *Chiroteuthis veranyi*, *Selenoteuthis scintillans*, *Clupea harengus*, and *Merluccius bilinearis* were found only as trace items and comprised a small part of the diet (Table 5).

DISCUSSION

The most striking aspect of these data was the strong dominance of *Loligo pealei* in every prey importance category. The relatively high prey diversity was also noteworthy. Of the ten prey species identified, *Clupea harengus*, *Mer-*

Table 7. Trace and non-trace indices of relative importance (IRI) and modified mass, with associated ranks (#), for the prey of long-finned pilot whales stranded along the U.S. mid-Atlantic coast.

Species	Trace		Non-trace		Mod. mass	#
	IRI	#	IRI	#		
<i>L. pealei</i>	0.92	1	1.09	1	47.4%	1
<i>S. acanthias</i>	0.01	5	0.11	2	30.6%	2
Ommastrephidae	0.18	2	—	—	7.4%	3
<i>C. harengus</i>	<0.01	6	—	—	4.7%	4
<i>S. scombrus</i>	<0.01	6	0.03	3	4.6%	5
<i>H. reversa</i>	0.13	3	—	—	4.3%	6
<i>C. veranyi</i>	0.05	4	—	—	<1%	7
<i>Urophycis</i> spp.	<0.01	6	—	—	—	—
<i>M. bilinearis</i>	<0.01	6	—	—	—	—

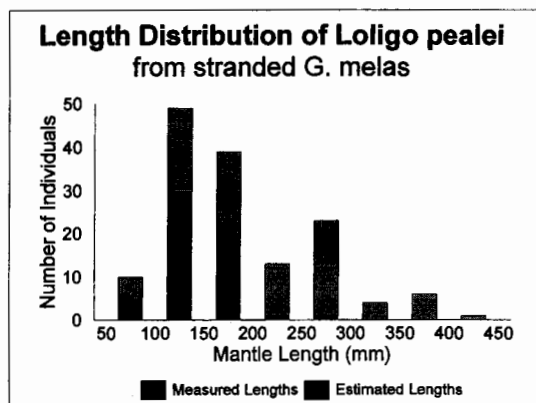


Figure 2. Length frequency distribution of *Loligo pealei* recovered from the stomach contents of stranded long-finned pilot whales.

luccius bilinearis, *Squalus acanthias*, *Urophycis* spp., *Chiroteuthis veranyi*, and *Selenoteuthis scintillans* have never before been reported in the diet of long-finned pilot whales. *Histioteuthis reversa* had previously only been known from pilot whales in the eastern North Atlantic (Desportes and Mouritsen 1993). Ten prey species were represented by trace specimens, while only three prey species had non-trace representatives. Previous studies in the Northwest Atlantic only analyzed intact, or non-trace, food material (Sergeant 1962, Mercer 1967, Mercer 1975, Waring *et al.* 1990, and Overholtz and Waring 1991), while studies in other regions utilized trace and non-trace remains (Martin *et al.* 1987, Gales and Pemberton 1992, Desportes and Mouritsen 1993). Our results indicate that the diet of western North Atlantic long-finned pilot whales differs substantially from that described previously, both in terms of diversity and relative species importance. Furthermore, the apparent difference in prey diversities noted between studies in the Northwest Atlantic and those in other regions seems to have been caused by differences in the treatment of trace and non-trace food remains.

Estimated dietary contributions of each prey species varied widely among the nine methods used to assess relative prey importance, indicating that it is difficult to make comparisons among food-habits studies that utilize different methodologies. Each method measures different parameters and each contains biases. For example, squid flesh has been shown to digest faster than fish flesh in the stomachs of marine mammals (Bigg and Fawcett 1985). Squid beaks, however, are resistant to breakdown by gastric activity and are known to persist in the stomachs of predators, while fish, including soft tissue and bones, are digestible (Bigg and Fawcett 1985). Therefore, analyses which take only intact items into account are likely to underestimate the importance of cephalopods. Studies which examine all stomach contents in one data set are liable to overestimate the importance of cephalopods. This is demonstrated in our data by the paucity of squid species represented by non-trace specimens. Since

the relative magnitudes of these biases are unknown, it is best to use a variety of trace and non-trace techniques in order to gain a more complete understanding of the diet.

Frequency-of-occurrence methods can overstate the importance of small or infrequently consumed prey, since neither prey size nor abundance is considered (Hyslop 1980, Bigg and Fawcett 1985, Bigg and Perez 1985, Pierce and Boyle 1991). Likewise, numerical abundance methods also exaggerate the contribution of small species. The concept behind IRI is that biases associated with the three component methods (FO, %Num, and %Mass) will be cancelled out by each other. However, because some biases afflict more than one component method, errors are probably compounded (Hyslop 1980). Of the methods used here, modified mass is perhaps the best single measure of relative prey importance, since it integrates non-trace frequency of occurrence, trace reconstructed mass, and non-trace reconstructed mass in a way that minimizes the effect of accumulated non-digestible parts (*i.e.*, squid beaks).

Because the lactation period is highly variable (Martin and Rothery 1993), body length is not a reliable indicator of nutritional independence for young pilot whales. However, we believed that it was important to have some knowledge about the extent to which our results may have been influenced by the inclusion of calves, which probably had a mixed diet of milk and solid food, and length was the only available criterion by which to make this judgment. We noted several gross differences between the stomach contents of the three calves and those of the older animals. The solid portion of the calf diet was generally characterized by: (1) small squids; (2) an absence of *Loligo pealei* (calf stomachs contained an average of 0.3 *L. pealei* beaks, while the entire sample averaged 79.4 *L. pealei* beaks per stomach); (3) few prey items (calf stomachs contained an average of 10 prey items compared to an average of 153 for the entire sample); (4) an absence of non-trace food remains; and (5) a complete lack of fish remains. Thus, it is likely that there are age-related dietary differences and that our results represent the combined diets of weaning calves and older, nutritionally independent animals. The prolonged weaning period is a learning stage during which calves probably experiment with solid food that is easy to capture. Smith and Read (1992) documented a transitional diet in weaning harbor porpoise (*Phocoena phocoena*) calves. Porpoise calves supplement milk obtained from their mothers with small, slow-moving euphausiids. By the time they are fully weaned, the diet of porpoise calves resembles that of adults and primarily consists of finfish (Read *et al.* 1994).

The limitations of food-habits data obtained from stranded marine mammals must be taken into account when interpreting these results. Our sample may have been biased toward nearshore prey (Clarke 1986b) or food that is not usually an important part of the diet. Illness, either caused by or resulting in malnutrition, can precipitate stranding. Nonetheless, the dietary composition of this sample is similar to that obtained from 30 pilot whales captured in the continental slope waters by the pelagic mackerel fishery (Gannon 1995). Based on a comparison of cetaceans collected at sea and from strandings, Ross

(1979) concluded that his sample of stranded animals provided representative dietary information.

Investigations that utilize small sample sizes, such as this, are vulnerable to biases. A skewed representation of the diet might be obtained if any animal had recently consumed prey that is not regularly eaten. This may account for the unexpectedly high level of importance given to the spiny dogfish. The large size difference between the dogfish and all other prey hints that dogfish might not be eaten frequently. The present sample included a very narrow segment of the population, collected predominantly during the spring. No adult males and no immature females were present. Therefore, potential dietary differences associated with sex, maturity status, and season could not be investigated. In the future, we hope to examine a larger sample with representatives from each age/sex class and each season, including animals obtained from mass strandings and incidental fishery captures.

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